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50 gaa agc aac att aca gtt ctc ata aag ctg ggg acc ccg act ctg ctg
438
Glu Ser Asn Ile Thr Val Leu Ile Lys Leu Gly Thr Pro Thr Leu Leu
40 45 50
55 gca aaa ccc tgt tac atc gtc att tct aaa aga cat ata acc atg ttg
486
Ala Lys Pro Cys Tyr Ile Val Ile Ser Lys Arg His Ile Thr Met Leu

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	ggc act gtg tcc cggtc aag atg caa gaa gga gtg aaa atg gcc tta 870			
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780

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Ala Leu Pro Arg Glu Ser Asn Ile Thr Val Leu Ile Lys Leu Gly Thr
35 40 45

35 Pro Thr Leu Leu Ala Lys Pro Cys Tyr Ile Val Ile Ser Lys Arg His
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Ile Thr Met Leu Ser Ile Lys Ser Gly Glu Arg Ile Val Phe Thr Phe
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Ser Cys Gln Ser Pro Glu Asn His Phe Val Ile Glu Ile Gln Lys Asn
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45 Ile Asp Cys Met Ser Gly Pro Cys Pro Phe Gly Glu Val Gln Leu Gln
100 105 110

Pro Ser Thr Ser Leu Leu Pro Thr Leu Asn Arg Thr Phe Ile Trp Asp
115 120 125

50 Val Lys Ala His Lys Ser Ile Gly Leu Glu Leu Gln Phe Ser Ile Pro
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Arg Leu Arg Gln Ile Gly Pro Gly Glu Ser Cys Pro Asp Gly Val Thr
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His Ser Ile Ser Gly Arg Ile Asp Ala Thr Val Val Arg Ile Gly Thr
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	Phe Cys Ser Asn Gly Thr Val Ser Arg Ile Lys Met Gln Glu Gly Val			
	180	185	190	
5	Lys Met Ala Leu His Leu Pro Trp Phe His Pro Arg Asn Val Ser Gly			
	195	200	205	
	Phe Ser Ile Ala Asn Arg Ser Ser Ile Lys Arg Leu Cys Ile Ile Glu			
	210	215	220	
10	Ser Val Phe Glu Gly Glu Gly Ser Ala Thr Leu Met Ser Ala Asn Tyr			
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15	Pro Glu Gly Phe Pro Glu Asp Glu Leu Met Thr Trp Gln Phe Val Val			
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	Pro Ala His Leu Arg Ala Ser Val Ser Phe Leu Asn Phe Asn Leu Ser			
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	Cys Thr Asp His Arg Tyr Cys Gln Arg Lys Ser Tyr Ser Leu Gln Val			
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	Lys Leu Leu Val Pro Lys Asp Arg Leu Ser Leu Val Leu Val Pro Ala			
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	Gln Lys Leu Gln Gln His Thr His Glu Lys Pro Cys Asn Thr Ser Phe			
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Ser Tyr Leu Val Ala Ser Ala Ile Pro Ser Gln Asp Leu Tyr Phe Gly
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5 Ser Phe Cys Pro Gly Gly Ser Ile Lys Gln Ile Gln Val Lys Gln Asn
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Asn Ile Ser Val Pro Arg Asp Gln Val Ala Cys Leu Thr Phe Phe Lys
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Glu Arg Ser Gly Val Val Cys Gln Thr Gly Arg Ala Phe Met Ile Ile
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Gln Glu Gln Arg Thr Arg Ala Glu Glu Ile Phe Ser Leu Asp Glu Asp
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Ser Asn Cys Ser Pro Thr Ser Gly Lys Gln Leu Asp Leu Leu Phe Ser
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35

Val Thr Leu Thr Pro Arg Thr Val Asp Leu Thr Val Ile Leu Ile Ala
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Cys Cys Val Lys Lys Lys Lys Lys Lys Thr Asn Lys Gly Pro Ala Val
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Lys Phe Gln Lys Gly Arg Lys Asp Asn Asp Ser His Val Tyr Ala Val
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Ile Glu Asp Thr Met Val Tyr Gly His Leu Leu Gln Asp Ser Ser Gly
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Met Gly Val Cys Pro Pro Ser Pro Pro Thr Ile Cys Ser Arg Ala Pro
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Thr Ala Lys Leu Ala Thr Glu Glu Pro Pro Pro Arg Ser Pro Pro Glu
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ttgaattcac tgtgtggagc c
21

5 <210> 29
 <211> 19
 <212> DNA
 <213> Artificial sequence

10 <220>
 <223> Description of the artificial sequence: Primer

 <400> 29
 tgcaggcaac agtgatgtc
 19
15

 <210> 30
 <211> 24
20 <212> DNA
 <213> Artificial sequence

 <220>
 <223> Description of the artificial sequence: Primer
25
 <400> 30
 attggccttc cttaggctgg ctac
 24

30

 <210> 31
 <211> 43
 <212> DNA
35 <213> Artificial sequence

 <220>
 <223> Description of the artificial sequence: Primer

40 <400> 31
 tgttagcgtga agacgacaga aagggcgtgg taccgagctc gag
 43

45
 <210> 32
 <211> 22
 <212> DNA
 <213> Artificial sequence
50
 <220>
 <223> Description of the artificial sequence: Primer

 <400> 32
55 agggcgtggt accgagctcg ag
 22

5 <210> 33
 <211> 11
 <212> DNA
 5 <213> Artificial sequence

 <220>
 <223> Description of the artificial sequence: Primer

10 <400> 33
 ggctcgagct c
 11

15 <210> 34
 <211> 22
 <212> DNA
 <213> Artificial sequence
20 <220>
 <223> Description of the artificial sequence: Primer

 <400> 34
25 ggccatgtcc ggtgggcttg tg
 22

30 <210> 35
 <211> 26
 <212> DNA
 <213> Artificial sequence

35 <220>
 <223> Description of the artificial sequence: Primer

 <400> 35
 ctcaaaaactc ctggacaagt tgctgg
40 26

 <210> 36
 <211> 22
45 <212> DNA
 <213> Artificial sequence

 <220>
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50 <400> 36
 aagggtgaagg tcggagtcaa cg
 22

55 <210> 37
 <211> 24

<212> DNA
<213> Artificial sequence

5 <220>
<223> Description of the artificial sequence: Primer

<400> 37
ggcagagatg atgacccttt tggc
24

10

<210> 38
<211> 23

15 <212> DNA
<213> Artificial sequence

<220>
<221> 5'UTR
20 <222> (1)..(282)

<220>
<221> GC_signal
<222> (147)..(157)

25 <220>
<221> misc_feature
<222> (201)..(209)
<223> cap signal; Transcription start

30 <220>
<221> 3'UTR
<222> (2794)..(6163)

35 <220>
<221> 3'UTR
<222> (2794)..(6163)

40 <220>
<221> CDS
<222> (283)..(2793)

<400> 38
agcagcgaaa ccccttagcag tgc

45 23

<210> 39
<211> 26

50 <212> DNA
<213> Artificial sequence

<220>
<223> Description of the artificial sequence: Primer

55 <400> 39
agaaccccta gcagtgcgtt agagac
26

5 <210> 40
5 <211> 27
5 <212> DNA
5 <213> Artificial sequence

10 <220>
10 <223> Description of the artificial sequence: Primer

15 <400> 40
15 gaactgtaat gttgcttct cgtggca
15 27